

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/535,522  
Source: PCT/10  
Date Processed by STIC: 5/27/05

***ENTERED***

Raw Sequence Listing before editing,  
for reference only



PCT

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/535,522

DATE: 05/27/2005  
TIME: 11:22:32

Input Set : A:\PTO.DA.txt  
Output Set: N:\CRF4\05272005\J535522.raw

3 <110> APPLICANT: Steinkasserer, Alexander  
5 <120> TITLE OF INVENTION: Use of Soluble Forms of CD83 and Nucleic Acids Encoding  
6 them for the Treatment or Prevention of Diseases  
W--> 7 <130> FILE REFERENCE: 032723woJH  
W--> 8 <140> CURRENT APPLICATION NUMBER:  
C--> 9 <141> CURRENT FILING DATE: 2005-05-18  
W--> 10 <160> NUMBER OF SEQ ID: 12  
11 <170> SOFTWARE: PatentIn Ver. 2.1

## ERRORED SEQUENCES

*Does Not Comply  
Corrected Diskette Needed*

356 <210> SEQ ID NO: 8  
357 <211> LENGTH: 139  
358 <212> TYPE: PRT  
359 <213> ORGANISM: Artificial Sequence *insert <220>*  
W--> 360 <220> FEATURE:  
360 <223> OTHER INFORMATION: Description of Artificial Sequence: partial  
361 sequence of pGEX2ThCD83ext  
OK> 363 <400> SEQUENCE: 8  
364 Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu  
365 -5 -1 1 5  
367 Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro  
368 10 15 20  
370 Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu  
371 25 30 35  
373 Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly  
374 40 45 50 55  
376 Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn  
377 60 65 70  
379 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser  
380 75 80 85  
382 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu  
383 90 95 100  
385 Ser Gly Lys Val Ile Leu Arg Val Thr Gly Cys Pro Ala Gln Arg Lys  
386 105 110 115  
388 Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile  
389 120 125 130  
444 <210> SEQ ID NO: 10  
445 <211> LENGTH: 139  
446 <212> TYPE: PRT  
447 <213> ORGANISM: Artificial Sequence  
W--> 448 <220> FEATURE:

*insert <220>*

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/535,522

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Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05272005\J535522.raw

448 <223> OTHER INFORMATION: Description of Artificial Sequence: partial  
449 sequence of pGEX2ThCD83ext\_mut129\_CtoS  
451 <400> SEQUENCE: 11 10  
452 Pro Pro Lys Ser Asp Leu Val Pro Arg Gly Ser Pro Gly Thr Pro Glu  
453 -5 -1 1 5  
455 Val Lys Val Ala Cys Ser Glu Asp Val Asp Leu Pro Cys Thr Ala Pro  
456 10 15 20  
458 Trp Asp Pro Gln Val Pro Tyr Thr Val Ser Trp Val Lys Leu Leu Glu  
459 25 30 35  
461 Gly Gly Glu Glu Arg Met Glu Thr Pro Gln Glu Asp His Leu Arg Gly  
462 40 45 50 55  
464 Gln His Tyr His Gln Lys Gly Gln Asn Gly Ser Phe Asp Ala Pro Asn  
465 60 65 70  
467 Glu Arg Pro Tyr Ser Leu Lys Ile Arg Asn Thr Thr Ser Cys Asn Ser  
468 75 80 85  
470 Gly Thr Tyr Arg Cys Thr Leu Gln Asp Pro Asp Gly Gln Arg Asn Leu  
471 90 95 100  
473 Ser Gly Lys Val Ile Leu Arg Val Thr Gly Ser Pro Ala Gln Arg Lys  
474 105 110 115  
476 Glu Glu Thr Phe Lys Lys Tyr Arg Ala Glu Ile  
478 120 125 130  
494 <210> SEQ ID NO: 12  
495 <211> LENGTH: 66  
496 <212> TYPE: DNA  
497 <213> ORGANISM: Artificial Sequence  
W--> 498 <220> FEATURE:  
499 <223> OTHER INFORMATION: Description of Artificial Sequence: primer  
500 antisense-CD83extra\_mutantCtoS  
502 <400> SEQUENCE: 12  
503 aattagaatt ctc当地atctc cgctctgtat ttcttaaaag tctttttt acgtgtgca 60  
504 gggat 66  
E--> 509 2 -

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/535,522

DATE: 05/27/2005

TIME: 11:22:33

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\05272005\J535522.raw

L:7 M:283 W: Missing Blank Line separator, <130> field identifier  
L:8 M:283 W: Missing Blank Line separator, <140> field identifier  
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:10 M:283 W: Missing Blank Line separator, <160> field identifier  
L:18 M:283 W: Missing Blank Line separator, <220> field identifier  
L:128 M:283 W: Missing Blank Line separator, <220> field identifier  
L:286 M:283 W: Missing Blank Line separator, <220> field identifier  
L:297 M:283 W: Missing Blank Line separator, <220> field identifier  
L:308 M:283 W: Missing Blank Line separator, <220> field identifier  
L:311 M:283 W: Missing Blank Line separator, <220> field identifier  
L:314 M:283 W: Missing Blank Line separator, <220> field identifier  
L:360 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:363 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:8  
L:396 M:283 W: Missing Blank Line separator, <220> field identifier  
L:399 M:283 W: Missing Blank Line separator, <220> field identifier  
L:402 M:283 W: Missing Blank Line separator, <220> field identifier  
L:448 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10  
L:451 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:10  
L:451 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:10 differs:11  
L:486 M:283 W: Missing Blank Line separator, <220> field identifier  
L:498 M:283 W: Missing Blank Line separator, <220> field identifier  
L:509 M:254 E: No. of Bases conflict, this line has no nucleotides.